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INFORMATION DISCLOSURE STATEMENT BY APPLICANT

(Use as many sheets as necessary)

Sheet 1 of 3

Complete if Known	
Application Number	~~Not yet assigned~~ 10/752,239
Filing Date	~~Herewith~~ 01/06/04
First Named Inventor	Steven D. Schwartz
Art Unit	~~Not yet assigned~~ 2121
Examiner Name	~~Not yet assigned~~ Joseph P. Hirl
Attorney Docket Number	96700/855

U. S. PATENT DOCUMENTS

Examiner Initials*	Cite No.[1]	Document Number Number-Kind Code[2] (if known)	Publication Date MM-DD-YYYY	Name of Patentee or Applicant of Cited Document	Pages, Columns, Lines, Where Relevant Passages or Relevant Figures Appear
JH	1	US- 6,185,548 B1	02-06-2001	Schwartz et al.	
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FOREIGN PATENT DOCUMENTS

Examiner Initials*	Cite No.[1]	Foreign Patent Document Country Code[3]-Number[4]-Kind Code[5] (if known)	Publication Date MM-DD-YYYY	Name of Patentee or Applicant of Cited Document	Pages, Columns, Lines, Where Relevant Passages Or Relevant Figures Appear	T[6]

Examiner Signature		Date Considered	1/5/05

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Application Number	~~Not yet assigned~~ 10/752,259
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Examiner Name	~~Not yet assigned~~ Joseph P. Hirl
Attorney Docket Number	96700/855

Sheet 2 of 3

NON PATENT LITERATURE DOCUMENTS

Examiner Initials*	Cite No.[1]	Include name of the author (in CAPITAL LETTERS), title of the article (when appropriate), title of the item (book, magazine, journal, serial, symposium, catalog, etc.), date, page(s), volume-issue number(s), publisher, city and/or country where published.	T[2]
H	2	BAGDASSARIAN et al., Molecular Electrostatic Potential Analysis for Enzymatic Substrates, Competitive Inhibitors, and Transition-State Inhibitors. J. Am. Chem. Soc., 118:8825-36, 1996.	
H	3	BETTS et al., Cytidine Deaminase. The 2-3 Angstrom Crystal Structure of an Enzyme: Transition-state Analog Complex. J. Mol. Biol., 235:635-56, 1994.	
H	4	BOHM, New Approaches in Molecular Structure Prediction. Biophysical Chemistry, 59:1-32, 1996.	
H	5	BRUSIC et al., Prediction of MHC Class II-Binding Peptides Using an Evolutionary Algorithm and Artificial Neural Network. Bioinformatics, 14:121-30, 1998.	
H	6	EHRLICH and SCHRAMM, Electrostatic Potential Surface Analysis of the Transition State for AMP Nucleosidase and for Formycin 5'-Phosphate, a Transition-State Inhibitor. Biochem., 33:8890-96, 1994.	
H	7	FRICK et al., Binding of Pyrimidin-2-one Ribonucleoside by Cytidine Deaminase as the Transition-State Analogue 3,4-Dihydrouridine and the Contribution of the 4-Hydroxyl Group to Its Binding Affinity. Biochemistry, 28:9423-30, 1989.	
H	8	GASTEIGER et al., Representation of Molecular Electrostatic Potentials by Topological Feature Maps. J. Am. Chem. Soc., 116:4608-20, 1994.	
H	9	HORENSTEIN and SCHRAMM, Electronic Nature of the Transition State for Nucleoside Hydrolase. A Blueprint for Inhibitor Design. Biochemistry, 32:7089-97, 1993.	
H	10	KLINE and SCHRAMM, Electrostatic Potential Surfaces of the Transition State for AMP Deaminase and for (R)-Coformycin, a Transition State Inhibitor. J. Biol Chem., 269:22385-90, 1994.	
H	11	SO and RICHARDS, Application of Neural Networks: Quantitative Structure-Activity Relationships of the Derivatives of 2,4-Diamino-5-(substituted-benzyl) pyrimidines as DHFR Inhibitors. J. Med. Chem., 35:3201-7, 1992.	

Examiner Signature		Date Considered	1/5/5

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JH	12	WAGENER et al., Autocorrelation of Molecular Surface Properties for Modeling Corticosteriod Binding Globulin and Cytosolic Ah Receptor Activity by Neural Networks. J. Am. Chem. Soc., 117:7769-75, 1995.	
JH	13	WEINSTEIN et al., Predictive Statistics and Artificial Intelligence in the U.S. National Cancer Institute's Drug Discovery Program for Cancer and AIDS. Stem Cells, 12:13-22, 1994.	

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